	ATGAAATTTA GTAAAAAATA TATAGCAGCT GGATCAGCTG TTATCGTATC CTTGAGTCTA	
	TGTGCCTATG CACTAAACCA GCATCGTTCG CAGGAAAATA AGGACAATAA TCGTGTCTCT	60
	TATGTGGATG GCAGCCAGTC AAGTCAGAAA AGTGAAAACT TGACACCAGA CCAGGTTAGC	120
	CAGAAAGAAG GAATTCAGGC TGAGCAAATT GTAATCAAAA TTACAGATCA GGGCTATGTA	180
5	ACGTCACACG GTGACCACTA TCATTACTAT AATGGGAAAG TTCCTTATGA TGCCCTCTTT	240
	AGTGAAGAAC TCTTGATGAA GGATCCAAAC TATCAACTTA AAGACGCTGA TATTGTCAAT	300
•	GAAGTCAAGG CTGGTTATAT CATCAAGGTC GATGGAAAAT ATTATGTCTA CCTGAAAGAT	360
	GCAGCTCATG CTGATAATGT TCGAACTAAA GATGAAATCA ATCGTCAAAA ACAAGAACAT	420
	CTCAAAGATA ATGAGAAGGT TAACTCTAAT GTTGCTGTAG CAAGGTCTCA GGGACGATAT	480
10	ACGACAAATG ATGGTTATGT CTTTAATCCA GCTGATATTA TCGAAGATAC GGGTAATGCT	540
	TATATCGTTC CTCATGGAGG TCACTATCAC TACATTCCCA AAAGCGATTT ATCTGCTAGT	600
	GAATTAGCAG CAGCTAAAGC ACATCTGGCT GGAAAAAATA TGCAACCGAG TCAGTTAAGC	660
	TATTCTTCAA CAGCTAGTGA CAATAACACG CAATCTGTAG CAAAAGGATC AACTAGCAAG	720
		780
15	CCAGCAAATA AATCTGAAAA TCTCCAGAGT CTTTTGAAGG AACTCTATGA TTCACCTAGC	840
15	GCCCAACGTT ACAGTGAATC AGATGGCCTG GTCTTTGACC CTGCTAAGAT TATCAGTCGT	900
	ACACCAAATG GAGTTGCGAT TCCGCATGGC GACCATTACC ACTTTATTCC TTACAGCAAG	960
	CTTTCTGCTT TAGAAGAAAA GATTGCCAGA ATGGTGCCTA TCAGTGGAAC TGGTTCTACA	1020
	GTTTCTACAA ATGCAAAACC TAATGAAGTA GTGTCTAGTC TAGGCAGTCT TTCAAGCAAT	1080
20	CCTTCTTCTT TAACGACAAG TAAGGAGCTC TCTTCAGCAT CTGATGGTTA TATTTTTAAT	1140
20	CCAAAAGATA TCGTTGAAGA AACGGCTACA GCTTATATTG TAAGACATGG TGATCATTTC	1200
	CATTACATTC CAAAATCAAA TCAAATTGGG CAACCGACTC TTCCAAACAA TAGTCTAGCA	1260
	ACACCTTCTC CATCTCTTCC AATCAATCCA GGAACTTCAC ATGAGAAACA TGAAGAAGAT	1320
	GGATACGGAT TTGATGCTAA TCGTATTATC GCTGAAGATG AATCAGGTTT TGTCATGAGT	1380
25	CACGGAGACC ACAATCATTA TTTCTTCAAG AAGGACTTGA CAGAAGAGCA AATTAAGGCT	1440
25	GCGCAAAAAC ATTTAGAGGA AGTTAAAACT AGTCATAATG GATTAGATTC TTTGTCATCT	1500
	CATGAACAGG ATTATCCAGG TAATGCCAAA GAAATGAAAG ATTTAGATAA AAAAATCGAA	1560
	GAAAAATTG CTGGCATTAT GAAACAATAT GGTGTCAAAC GTGAAAGTAT TGTCGTGAAT	1620
	AAAGAAAAA ATGCGATTAT TTATCCGCAT GGAGATCACC ATCATGCAGA TCCGATTGAT	1680
20	GAACATAAAC CGGTTGGAAT TGGTCATTCT CACAGTAACT ATGAACTGTT TAAACCCGAA	1740
30	CAAGGAGTTG CTAAAAAAAGA AGGGAATAAA GTTTATACTG GAGAAGAATT AACGAATGTT	1800
	GTTAATTTGT TAAAAAATAG TACGTTTAAT AATCAAAACT TTACTCTAGC CAATGGTCAA	1860
	AAACGCGTTT CTTTTAGTTT TCCGCCTGAA TTGGAGAAAA AATTAGGTAT CAATATGCTA	1920
	GTAAAATTAA TAACACCAGA TGGAAAAGTA TTGGAGAAAG TATCTGGTAA AGTATTTGGA	1980
	GAAGGAGTAG GGAATATTGC AAACTTTGAA TTAGATCAAC CTTATTTACC AGGACAAACA	2040
35	TTTAAGTATA CTATCGCTTC AAAAGATTAT CCAGAAGTAA GTTATGATGG TACATTTACA	2100
	GTTCCAACCT CTTTAGCTTA CAAAATGGCC AGTCAAACGA TTTTCTATCC TTTCCATGCA	2160
	GGGGATACTT ATTTAAGAGT GAACCCTCAA TTTGCAGTGC CTAAAGGAAC TGATGCTTTA	2220
	GTCAGAGTGT TTGATGAATT TCATGGAAAT GCTTATTTAG AAAATAACTA TAAAGTTGGT	2280
	GAAATCAAAT TACCGATTCC GAAATTAAAC CAAGGAACAA CCAGAACGGC CGGAAATAAA	2340
40	ATTCCTGTAA CCTTCATGGC AAATGCTTAT TTGGACAATC AATCGACTTA TATTGTGGAA	2400
	GTACCTATCT TGGAAAAAGA AAATCAAACT GATAAACCAA GTATTCTACC ACAATTTAAA	2460
	AGGAATAAAG CACAAGAAAA CTCAAAACTT GATGAAAAGG TAGAAGAACC AAAGACTAGT	2520
	GAGAAGGTAG AAAAAGAAAA ACTTTCTGAA ACTGGGAATA GTACTAGTAA TTCAACGTTA	2580
	GAAGAAGTTC CTACAGTGGA TCCTGTACAA GAAAAAGTAG CAAAATTTGC TGAAAGTTAT	2640
45	GGGATGAAGC TAGAAAATGT CTTGTTTAAT ATGGACGGAA CAATTGAATT ATATTTACCA	2700

	TCAGGAGAAG TCATTAAAAA GAATATGGCA GATTTTACAG GAGAAGCACC TCAAGGAAAT	2760
	GGTGAAAATA AACCATCTGA AAATGGAAAA GTATCTACTG GAACAGTTGA GAACCAACCA	2820
	ACAGAAAATA AACCAGCAGA TTCTTTACCA GAGGCACCAA ACGAAAAACC TGTAAAACCA	2880
	GAAAACTCAA CGGATAATGG AATGTTGAAT CCAGAAGGGA ATGTGGGGAG TGACCCTATG	2940
5	TTAGATCCAG CATTAGAGGA AGCTCCAGCA GTAGATCCTG TACAAGAAAA ATTAGAAAAA	3000
	TTTACAGCTA GTTACGGATT AGGCTTAGAT AGTGTTATAT TCAATATGGA TGGAACGATT	3060
	GAATTAAGAT TGCCAAGTGG AGAAGTGATA AAAAAGAATT TATCTGATTT CATAGCGTAA	3120
	(SEQ ID NO: 1)	
	FIGURE 1	
10		
	AATTCCTTGT CGGGTAAGTT CCGACCCGCA CGAAAGGCGT AATGATTTGG GCACTGTCTC	60
	AACGAGAGAC TCGGTGAAAT TTTAGTACCT GTGAAGATGC AGGTTACCCG CGACAGGACG	120
	GAAAGACCCC ATGGAGCTTT ACTGCAGTTT GATATTGAGT GTCTGTACCA CATGTACAGG	180
15	ATAGGTAGGA GTCTAAGAGA TCGGGACGCC AGTTTCGAAG GAGACGCTGT TGGGATACTA	240
	CCCTTGTGTT ATGGCCACTC TAACCCAGAT AGGTGATCCC TATCGGAGAC AGTGTCTGAC	300
	GGGCAGTTTG ACTGGGGCGG TCGCCTCCTA AAAGGTAACG GAGGCGCCCA AAGGTTCCCT	360
	CAGAATGGTT GGAAATCATT CGCAGAGTGT AAAGGTATAA GGGAGCTTGA CTGCGAGAGC	420
	TACAACTCGA GCAGGGACGA AAGTCGGGCT TAGTGATCCG GTGGTTCCGT ATGGAAGGGC	480
20	CATCGCTCAA CGGATAAAAG CTACCCTGGG GATAACAGGC TTATCTCCCC CAAGAGTTCA	540
	CATCGACGGG GAGGTTTGGC ACCTCGATGT CGGCTCGTCG CATCCTGGGG CTGTAGTCGG	600
	TCCCAAGGGT TGGGCTGTTC GCCCATTAAA GCGGCACGCG AGCTGGGTTC AGAACGTCGT	660
	GAGACAGTTC GGTCCCTATC CGTCGCGGGC GTAGGAAATT TGAGAGGATC TCCTCCTAGT	720
	ACGAGAGGAC CAGAGTGGAC TTACCGCTGG TGTACCAGTT GTCTTGCCAA AGGCATCGCT	780
25	GGGTAGCTAT GTAGGGAAGG GATAAACGCT GAAAGCATCT AAGTGTGAAA CCCACCTCAA	840
	CATGAGATTT CCCATGATTA TATATCAGTA AGAGCCCTGA GAGATGATCA GGTAGATAGG	900
٠	TTAGAAGTGG AAGTGTGGCG ACACATGTAG CGGACTAATA CTAATAGCTC GAGGACTTAT	960
	CCAAAGTAAC TGAGAATATG AAAGCGAACG GTTTTCTTAA ATTGAATAGA TATTCAATTT	1020
	TGAGTAGGTA TTACTCAGAG TTAAGTGACG ATAGCCTAGG AGATACACCT GTACCCATGC	1080
30	CGAACACAGA AGTTAAGCCC TAGAACGCCG GAAGTAGTTG GGGGTTGCCC CCTGTGAGAT	1140
	AGGGAAGTCG CTTAGCTCTA GGGAGTTTAG CTCAGCTGGG AGAGCATCTG CCTTACAAGC	1200
	AGAGGGTCAG CGGTTCGATC CCGTTAACTC CCAAAGGTCC CGTAGTGTAG CGGTTATCAC	1260
	GTCGCCCTGT CACGGCGAAG ATCGCGGGTT CGATTCCCGT CGGGACCGTT TAAGGTAACG	1320
	CAAGTTATTT TAGACTCGTT AGCTCAGTTG GTAGAGCAAT TGACTTTTAA TCAATGGGTC	1380
35	ACTGGTTCGA GCCCAGTACG GGTCATATAT GCGGGTTTGG CGGAATTCTA ATCTCTTTGA	1440
	AATCATCTTC TCTCACTTTC CAAAACTCTA TTACCTCTTA TTATACCACA TTTCAATCTT	1500
	CAACTTCCCA GTAATATAAG CACCTCTGGC GAAAGAAGTT TCAATGTCCT AAAGTAATAA	1560
-	GTGAATCCAA TTCAGGAACT CCAAGAACAA AAGAAACATC TGGTGTCACA AGTATTGGAT	1620
40	GGCACAGAGT CACGTGGTAG TCTGACCCTA GCAGAAATTT TAAATAGTAA ACTATTTACT	1680
40	GGTTAATTAA ATGGTTAAAT AACCGGTTTA GAAAACTATT TAATAAAGTA AAAGAAGTTG	1740
	AGAAAAAACT TCATCATTTA TTGAAATGAG GGATTTATGA AATTTAGTAA AAAATATATA	1800
	GCAGCTGGAT CAGCTGTTAT CGTATCCTTG AGTCTATGTG CCTATGCACT AAACCAGCAT	1860
	CGTTCGCAGG AAAATAAGGA CAATAATCGT GTCTCTTATG TGGATGGCAG CCAGTCAAGT	1920
4 6	CAGAAAAGTG AAAACTTGAC ACCAGACCAG GTTAGCCAGA AAGAAGGAAT TCAGGCTGAG	1980
45	CAAATTGTAA TCAAAATTAC AGATCAGGGC TATGTAACGT CACACGGTGA CCACTATCAT	2040

	TACTATAATG GGAAAGTTCC TTATGATGCC CTCTTTAGTG AAGAACTCTT GATGAAGGAT	•
	CCAAACTATC AACTTAAAGA CGCTGATATT GTCAATGAAG TCAAGGGTGG TTATATCATC	2100
	AAGGTCGATG GAAAATATTA TGTCTACCTG AAAGATGCAG CTCATGCTGA TAATGTTCGA	2160
	ACTARAGATG ARATCARTCG TCARARACAR GARCATGTCA ARGATRATGA GARGGTTRAC	2220
5	TCTAATGTTG CTGTAGCAAG GTCTCAGGGA CGATATACGA CAAATGATGG TTATGTCTTT	2280
_	AATCCAGCTG ATATTATCGA AGATACGGGT AATGCTTATA TCGTTCCTCA TGGAGGTCAC	2340
	TATCACTACA TICCCAAAAG CGATTTATCT GCTAGTGAAT TAGCAGCAGC TAAAGCACAT	2400
	CTGGCTGGAA AAAATATGCA ACCGAGTCAG TTAAGCTATT CTTCAACAGC TAGTGACAAT	2460
	AACACGCAAT CTGTAGCAAA AGGATCAACT AGCAAGCCAG CAAATAAATC TGAAAATCTC	2520
10	CAGAGTCTTT TGAAGGAACT CTATGATTCA CCTAGCGCCC AACGTTACAG TGAATCAGAT	2580
	GCCCTGGTCT TTGACCCTGC TAAGATTATC AGTCGTACAC CAAATGGAGT TGCGATTCCG	2640
	CATGGCGACC ATTACCACTT TATTCCTTAC AGCAAGCTTT CTGCTTTAGA AGAAAAGATT	2700
	GCCAGAATGG TGCCTATCAG TGGAACTGGT TCTACAGTTT CTACAAATGC AAAACCTAAT	2760
	GAAGTAGTGT CTAGTCTAGG CAGTCTTTCA AGCAATCCTT CTTCTTTAAC GACAAGTAAG	2820
15	GAGCTCTCTT CAGCATCTGA TGGTTATATT TTTAATCCAA AAGATATCGT TGAAGAAACG	2880
	GCTACAGCTT ATATTGTAAG ACATGGTGAT CATTTCCATT ACATTCCAAA ATCAAATCAA	2940
		3000
	ATTGGGCAAC CGACTCTTCC AAACAATAGT CTAGCAACAC CTTCTCCATC TCTTCCAATC	3060
	AATCCAGGAA CTTCACATGA GAAACATGAA GAAGATGGAT ACGGATTTGA TGCTAATCGT	3120
20	ATTATCGCTG AAGATGAATC AGGTTTTGTC ATGAGTCACG GAGACCACAA TCATTATTTC	3180
	TTCAAGAAGG ACTTGACAGA AGAGCAAATT AAGGCTGCGC AAAAACATTT AGAGGAAGTT	3240
	AAAACTAGTC ATAATGGATT AGATTCTTTG TCATCTCATG AACAGGATTA TCCAGGTAAT	3300
	GCCAAAGAAA TGAAAGATTT AGATAAAAAA ATCGAAGAAA AAATTGCTGG CATTATGAAA	3360
	CAATATGGTG TCAAACGTGA AAGTATTGTC GTGAATAAAG AAAAAAATGC GATTATTTAT	3420
25	CCGCATGGAG ATCACCATCA TGCAGATCCG ATTGATGAAC ATAAACCGGT TGGAATTGGT	3480
23	CATTCTCACA GTAACTATGA ACTGTTTAAA CCCGAAGAAG GAGTTGCTAA AAAAGAAGGG	3540
	AATAAAGTTT ATACTGGAGA AGAATTAACG AATGTTGTTA ATTTGTTAAA AAATAGTACG	3600
	TTTAATAATC AAAACTTTAC TCTAGCCAAT GGTCAAAAAC GCGTTTCTTT TAGTTTTCCG	3660
	CCTGAATTGG AGAAAAAATT AGGTATCAAT ATGCTAGTAA AATTAATAAC ACCAGATGGA	3720
30	AAAGTATTGG AGAAAGTATC TGGTAAAGTA TTTGGAGAAG GAGTAGGGAA TATTGCAAAC	3780
20	TTTGAATTAG ATCAACCTTA TTTACCAGGA CAAACATTTA AGTATACTAT CGCTTCAAAA	3840
	GATTATCCAG AAGTAAGTTA TGATGGTACA TTTACAGTTC CAACCTCTTT AGCTTACAAA	3900
	ATGGCCAGTC AAACGATTTT CTATCCTTTC CATGCAGGGG ATACTTATTT AAGAGTGAAC	3960
	CCTCAATTG CAGTGCCTAA AGGAACTGAT GCTTTAGTCA GAGTGTTTGA TGAATTTCAT	4020
35	GGAAATGCTT ATTTAGAAAA TAACTATAAA GTTGGTGAAA TCAAATTACC GATTCCGAAA	4080
33	TTAAACCAAG GAACAACCAG AACGGCCGGA AATAAAATTC CTGTAACCTT CATGGCAAAT	4140
	GCTTATTTGG ACAATCAATC GACTTATATT GTGGAAGTAC CTATCTTGGA AAAAGAAAAT	4200
	CAAACTGATA AACCAAGTAT TCTACCACAA TTTAAAAGGA ATAAAGCACA AGAAAACTCA	4260
	AAACTTGATG AAAAGGTAGA AGAACCAAAG ACTAGTGAGA AGGTAGAAAA AGAAAAACTT	4320
<i>4</i> ∩		4380
40	GTACAAGAAA AAGTAGCAAA ATTTGCTGAA AGTTATGGGA TGAAGCTAGA AAATGTCTTG	4440
	TTTAATATGG ACGGAACAAT TGAATTATAT TTACCATCAG GAGAAGTCAT TAAAAAGAAT	4500
	ATGGCAGATT TTACAGGAGA AGCACCTCAA GGAAATGGTG AAAATAAACC ATCTGAAAAT	4560
	GGAAAAGTAT CTACTGGAAC AGTTGAGAAC CAACCAACAG AAAATAAACC AGCAGATTCT	4620
45	TTACCAGAGG CACCAAACGA AAAACCTGTA AAACCAGAAA ACTCAACGGA TAATGGAATG	4680
45	TTGAATCCAG AAGGGAATGT GGGGAGTGAC CCTATGTTAG ATCCAGCATT AGAGGAAGCT	4740

ri e		
	CCAGCAGTAG ATCCTGTACA AGAAAAATTA GAAAAATTTA CAGCTAGTTA CGGATTAGGC	4000
	THOM ING THATATTCAA TATGGATGGA ACGATTGAAT TAACATTCCC AACTCCACO	4800 4860
	GIGAIAAAAA AGAATTTATC TGATTTCATA GCGTAAGGAA TAGCACTAGA AAAACTTCATA	4920
	AICHAAAATG AAGTTCTCTC AAAAGTTAGA AATAAAACTC TGACTTTCCC AGAATTTCAT	4980
	TTTATTATTA ATATATAAAA TTTCTTGACA TACAACTTAA AAAGAGGTGG AATATTTACT	
	AGITAATT (SEQ ID NO : 2)	5040 5048
	FIGURE 2	2048
		•
10	ATGAAAATCA ATAAAAAATA TCTAGCTGGG TCAGTAGCTA CACTTGTTTT AAGTGTCTGT	•
	GCTTATGAAC TAGGTTTGCA TCAAGCTCAA ACTGTAAAAG AAAATAATCG TGTTTCCTAT	60
	ATAGATGGAA AACAAGCGAC GCAAAAAACG GAGAATTTGA CTCCTGATGA GGTTAGCAAG	120
	CGTGAAGGAA TCAACGCCGA ACAAATCGTC ATCAAGATTA CGGATCAAGG TTATGTGACC	180
	TCTCATGGAG ACCATTATCA TTACTATAAT GGCAAGGTCC CTTATGATGC CATCATCAGT	240
15	GAAGAGCTCC TCATGAAAGA TCCGAATTAT CAGTTGAAGG ATTCAGACAT TGTCAATGAA	300
	ATCAAGGGTG GTTATGTCAT TAAGGTAAAC GGTAAATACT ATGTTTACCT TAAGGATGCA	360
	GCTCATGCGG ATAATGTCCG TACAAAAGAA GAAATCAATC GGCAAAAACA AGAACATAGT	420
	CAGCATCGTG AAGGAGGCAC TTGAGGAAAA GAAATCAATC GGCAAAAACA AGAACATAGT	480
	CAGCATCGTG AAGGAGGGAC TTCAGCAAAC GATGGTGCGG TAGCCTTTGC ACGTTCACAG	540
20	GGACGCTACA CCACAGATGA TGGTTATATC TTCAATGCAT CTGATATCAT CGAAGATACG	600
	GGCGATGCCT ATATCGTTCC TCATGGAGAT CATTACCATT ACATTCCTAA GAATGAGTTA	660
	TCAGCTAGCG AGTTGGCTGC TGCAGAAGCC TTCCTATCTG GTCGGGAAAA TCTGTCAAAT	720
	TTAAGAACCT ATCGCCGACA AAATAGCGAT AACACTCCAA GAACAAACTG GGTACCTTCT	780
	GTAAGCAATC CAGGAACTAC AAATACTAAC ACAAGCAACA ACAGCAACAC TAACAGTCAA	840
25	GCAAGTCAAA GTAATGACAT TGATAGTCTC TTGAAACAGC TCTACAAACT GCCTTTGAGT	900
23	CAACGCCATG TAGAATCTGA TGGCCTTATT TTCGACCCAG CGCAAATCAC AAGTCGAACC	960
	GCCAGAGGTG TAGCTGTCCC TCATGGTAAC CATTACCACT TTATCCCTTA TGAACAAATG	1020
	TCTGAATTGG AAAAACGAAT TGCTCGTATT ATTCCCCTTC GTTATCGTTC AAACCATTGG	1080
	GTACCAGATT CAAGACCAGA AGAACCAAGT CCACAACCGA CTCCAGAACC TAGTCCAAGT	1140
30	CCGCAACCTG CACCAAATCC TCAACCAGCT CCAAGCAATC CAATTGATGA GAAATTGGTC	1200
50	AAAGAAGCTG TTCGAAAAGT AGGCGATGGT TATGTCTTTG AGGAGAATGG AGTTTCTCGT	1260
	TATATCCCAG CCAAGAATCT TTCAGCAGAA ACAGCAGCAG GCATTGATAG CAAACTGGCC	1320
	AAGCAGGAAA GTTTATCTCA TAAGCTAGGA GCTAAGAAAA CTGACCTCCC ATCTAGTGAT	1380
	CGAGAATTTT ACAATAAGGC TTATGACTTA CTAGCAAGAA TTCACCAAGA TTTACTTGAT	1440
35	AATAAAGGTC GACAAGTTGA TTTTGAGGCT TTGGATAACC TGTTGGAACG ACTCAAGGAT	1500
33	GTCTCAAGTG ATAAAGTCAA GTTAGTGGAT GATATTCTTG CCTTCTTAGC TCCGATTCGT	1560
	CATCCAGAAC GTTTAGGAAA ACCAAATGCG CAAATTACCT ACACTGATGA TGAGATTCAA	1620
	GTAGCCAAGT TGGCAGGCAA GTACACAACA GAAGACGGTT ATATCTTTGA TCCTCGTGAT	1680
	ATAACCAGTG ATGAGGGGGA TGCCTATGTA ACTCCACATA TGACCCATAG CCACTGGATT	1740
40	AAAAAAGATA GTTTGTCTGA AGCTGAGAGA GCGGCAGCCC AGGCTTATGC TAAAGAGAAA	1800
40	GGTTTGACCC CTCCTTCGAC AGACCATCAG GATTCAGGAA ATACTGAGGC AAAAGGAGCA	1860
	GAAGCTATCT ACAACCGCGT GAAAGCAGCT AAGAAGGTGC CACTTGATCG TATGCCTTAC	1920
	AATCTTCAAT ATACTGTAGA AGTCAAAAAC GGTAGTTTAA TCATACCTCA TTATGACCAT	1980
	TACCATAACA TCAAATTTGA GTGGTTTGAC GAAGGCCTTT ATGAGGCACC TAAGGGGTAT	2040
4 ~	ACTOTTGAGG ATCTTTTGGC GACTGTCAAG TACTATGTCG AACATCCAAA CGAACGTCCG	2100
45	CATTCAGATA ATGGTTTTGG TAACGCTAGC GACCATGTTC AAAGAAACAA AAATGGTCAA	2160
		· •

•	GAACAAACGCA AAAACCAAGC GAGGAGAAAC CTCAGACAGA AAAACCTGAG	2220
	GAAGAAACCC CTCGAGAAGA GAAACCACAA AGCGAGAAAC CAGAGTCTCC AAAACCAACA	2280
	GAGGARCIAG AAGAAGAATC ACCAGAGGAA TCAGAAGAAC CTCAGGTCCA GAGRACAAC	2340
	OTTORAGARA AACTGAGAGA GGCTGAAGAT TTACTTGGAA AAATCCAGGA GCGAAGGAA	
	ANGICCAATG CCAAAGAGAC TCTCACAGGA TTAAAAAATA ATTTACTATT TCCCACAGGA	
-	GACAACAATA CTATTATGGC AGAAGCTGAA AAACTATTGG CTTTATTAAA GGACACTAAC	2520
	TAK (SEQ ID NO: 3)	2523
	FIGURE 3	4323
10		
	CAGAGATCTT AGTGAATCAA ATATACTTAA GAAAAGAGGA AAGAATGAAA ATCAATAAAA	60
	MATATCTAGC TGGGTCAGTA GCTACACTTG TTTTAAGTGT CTGTGCTTAT GAACTAGGTT	
	TGCATCAAGC TCAAACTGTA AAAGAAAATA ATCGTGTTTC CTATATAGAT CCAAAACAAC	180
	CGACGCAAAA AACGGAGAAT TTGACTCCTG ATGAGGTTAG CAAGCGTGAA GGAATCAACG	240
15	CCGAACAAT CGTCATCAAG ATTACGGATC AAGGTTATGT GACCTCTCAT GGAGACCATT	300
	ATCATTACTA TAATGGCAAG GTCCCTTATG ATGCCATCAT CAGTGAAGAG CTCCTCATGA	360
	AAGATCCGAA TTATCAGTTG AAGGATTCAG ACATTGTCAA TGAAATCAAG GGTGGTTATG	
	TCATTAAGGT AAACGGTAAA TACTATGTTT ACCTTAAGGA TGCAGCTCAT GCGGATAATG	420 480
	TCCGTACAAA AGAAGAAATC AATCGGCAAA AACAAGAACA TAGTCAGCAT CGTCAACCAC	•
20	GGACTTCAGC AAACGATGGT GCGGTAGCCT TTGCACGTTC ACAGGGACGC TACACCACAG	540
	ATGATGGTTA TATCTTCAAT GCATCTGATA TCATCGAAGA TACGGGCGAT GCCTATATCG	
	TTCCTCATGG AGATCATTAC CATTACATTC CTAAGAATGA GTTATCAGCT AGCGAGTTGG	660 720
	CTGCTGCAGA AGCCTTCCTA TCTGGTCGGG AAAATCTGTC AAATTTAAGA ACCTATCGCC	
	GACAAAATAG CGATAACACT CCAAGAACAA ACTGGGTACC TTCTGTAAGC AATCCAGGAA	780
25	CTACAAATAC TAACACAAGC AACAACAGCA ACACTAACAG TCAAGCAAGT CAAAGTAATG	840
	ACATTGATAG TCTCTTGAAA CAGCTCTACA AACTGCCTTT GAGTCAACGC CATGTAGAAT	
	CTGATGGCCT TATTTTCGAC CCAGCGCAAA TCACAAGTCG AACCGCCAGA GGTGTAGCTG	960
	TCCCTCATGG TAACCATTAC CACTTTATCC CTTATGAACA AATGTCTGAA TTGGAAAAAC	1020
	GAATTGCTCG TATTATTCCC CTTCGTTATC GTTCAAACCA TTGGGTACCA GATTCAAGAC	1080
30	CAGAAGAACC AAGTCCACAA CCGACTCCAG AACCTAGTCC AAGTCCGCAA CCTGCACCAA	1140
	ATCCTCAACC AGCTCCAAGC AATCCAATTG ATGAGAAATT GGTCAAAGAA GCTGTTCGAA	1200
	AAGTAGGCGA TGGTTATGTC TTTGAGGAGA ATGGAGTTTC TCGTTATATC CCAGCCAAGA	1260
	ATCTTTCAGC AGAAACAGCA GCAGGCATTG ATAGCAAACT GGCCAAGCAG GAAAGTTTAT	1320
	CTCATAAGCT AGGAGCTAAG AAAACTGACC TCCCATCTAG TGATCGAGAA TTTTACAATA	1380
35	AGGCTTATGA CTTACTAGCA AGAATTCACC AAGATTTACT TGATAATAAA GGTCGACAAG	1440
	TTGATTTTGA GGCTTTGGAT AACCTGTTGG AACGACTCAA GGATGTCTCA AGTGATAAAG	1500
	TCAAGTTAGT GGATGATATT CTTGCCTTCT TAGCTCCGAT TCGTCATCCA GAACGTTTAG	1560
	GAAAACCAAA TGCGCAAATT ACCTACACTG ATGATGAGAT TCAAGTAGCC AAGTTGGCAG	1620
	GCAAGTACAC AACAGAAGAC GGTTATATCT TTGATCCTCG TGATATAACC AGTGATGAGG	1680
40	GGGATGCCTA TGTAACTCCA CATATGACCC ATAGCCACTG GATTAAAAAA GATAGTTTGT	1740
	CTGAAGCTGA GAGAGCGGCA GCCCAGGCTT ATGCTAAAGA GAAAGGTTTG ACCCCTCCTT	1800
	CGACAGACCA TCAGGATTCA GGAAATACTG AGGCAAAAGG AGCAGAAGCT ATCTACAACC	1860
	GCGTGAAAGC AGCTAAGAAG GTGCCACTTG ATCGTATGCC TTACAATCTT CAATATACTG	1920
	TAGAAGTCAA AAACGGTAGT TTAATCATAC CTCATTATGA CCATTACCAT AACATCAAAT	1980
45	TTGAGTGGTT TGACGAAGGC CTTTATGAGG CACCTAAGGG GTATACTCTT GAGGATCTTT	2040
	GAGGATCTTT	2100

•	. The second contract \cdot	
	TEGERAL CAAGTACTAT GTCGAACATC CAAACGAACG TCCGCATTCA GATAATGGTT	24.52
	TIGGTAACGC TAGCGACCAT GTTCAAAGAA ACAAAAATGG TCAAGCTGAT ACCAAGCARA	2160
	CGGAAAAACC AAGCGAGGAG AAACCTCAGA CAGAAAAACC TGAGGAAGAA ACCCCTCCAG	2220
	AAGAGAAACC ACAAAGCGAG AAACCAGAGT CTCCAAAACC AACAGAGGAA CCAGAACAAC	2280
-	AATCACCAGA GGAATCAGAA GAACCTCAGG TCGAGACTGA AAAGGTTGAA GAAAAACTGA	2340
	GAGAGGCTGA AGATTTACTT GGAAAAATCC AGGATCCAAT TATCAAGTCC AATGCCAAAG	2400
	AGACTCTCAC AGGATTAAAA AATAATTTAC TATTTGGCAC CCAGGACAAC AATACTATTA	2460
	TGGCAGAAGC TGAAAAACTA TTGGCTTTAT TAAAGGAGAG TAAGTAAAGG TAGCAGCATT	2520
	TTCTAACTCC TAAAAACAGG ATAGGAGAAC GGGAAAACGA AAAATGAGAG CAGAATGTGA	2580
10	GTTCTAG (SED ID NO : 4)	2640
	FIGURE 4	2647
	GGGTCTTAAA ACTCTGAATC CTTTAGAGGC AGACCCACAA AATGACAAGA CCTATTTAGA	
15	AAATCTGGAA GAAAATATGA GTGTTCTAGC AGAAGAATTA AAGTGAGGAA AGAATGAAAA	60
	TCAATAAAAA ATATCTAGCA GGTTCAGTGG CAGTCCTTGC CCTAAGTGTT TGTTCCTATG	120
	AACTTGGTCG TCACCAAGCT GGTCAGGTTA AGAAAGAGTC TAATCGAGTT TCTTATATAG	180
	ATGGTGATCA GGCTGGTCAA AAGGCAGAAA ATTTGACACC AGATGAAGTC AGTAAGAGAG	240
	AGGGGATCAA CGCCGAACAA ATTGTTATCA AGATTACGGA TCAAGGTTAT GTGACCTCTC	300
20	ATGGAGACCA TTATCATTAC TATAATGGCA AGGTTCCTTA TGATGCCATC ATCAGTGAAG	360
	AACTTCTCAT GAAAGATCCG AATTATCAGT TGAAGGATTC AGACATTGTC AATGAAATCA	420
	AGGGTGGCTA TGTGATTAAG GTAGACGGAA AATACTATGT TTACCTTAAA GATGCGGCCC	480
	ATGCGGACAA TATTCGGACA AAAGAAGAGA TTAAACGTCA GAAGCAGGAA CACAGTCATA	540
	ATCATAACTC AAGAGCAGAT AATGCTGTTG CTGCAGCCAG AGCCCAAGGA CGTTATACAA	600
25	CGGATGATGG GTATATCTTC AATGCATCTG ATATCATTGA GGACACGGGT GATGCTTATA	660
	TCGTTCCTCA CGGCGACCAT TACCATTACA TTCCTAAGAA TGAGTTATCA GCTAGCGAGT	720
	TAGCTGCTGC AGAAGCCTAT TGGAATGGGA AGCAGGGATC TCGTCCTTCT TCAAGTTCTA	780
	GTTATAATGC AAATCCAGTT CAACCAAGAT TGTCAGAGAA CCACAATCTG ACTGTCACTC	840
	CAACTTATCA TCAAAATCAA GGGGAAAACA TTTCAAGCCT TTTACGTGAA TTGTATGCTA	900
30	AACCCTTATC AGAACGCCAT GTAGAATCTG ATGGCCTTAT TTTCGACCCA GCGCAAATCA	960
	CAAGTCGAAC CGCCAGAGGT GTAGCTGTCC CTCATGGTAA CCATTACCAC TTTATCCCTT	1020
	ATGAACAAAT GTCTGAATTG GAAAAACGAA TTGCTCGTAT TATTCCCCTT CGTTATCGTT	1080
	CAAACCATTG GGTACCAGAT TCAAGACCAG AACAACCAAG TCCACAATCG ACTCCGGAAC	1140
	CTAGTCCAAG TCTGCAACCT GCACCAAATC CTCAACCAGC TCCAAGCAAT CCAATTGATG	1200
35	AGAAATTGGT CAAAGAAGCT GTTCGAAAAG TAGGCGATGG TTATGTCTTT GAGGAGAATG	1260
	GAGTTTCTCG TTATATCCCA GCCAAGGATC TTTCAGCAGA AACAGCAGCA GGCATTGATA	1320
	GCAAACTGGC CAAGCAGGAA AGTTTATCTC ATAAGCTAGG AGCTAAGAAA ACTGACCTCC	1380
	CATCTAGTGA TCGAGAATTT TACAATAAGG CTTATGACTT ACTAGCAAGA ATTCACCAAG	1440
	ATTTACTTGA TAATAAAGGT CGACAAGTTG ATTTTGAGGT TTTGGATAAC CTGTTGGAAC	1500
40	GACTCAAGGA TGTCTCAAGT GATAAAGTCA AGTTAGTGGA TGATATTCTT GCCTTCTTAG	1560
	CTCCGATTCG TCATCCAGAA CGTTTAGGAA AACCAAATGC GCAAATTACC TACACTGATG	1620
	ATGAGATTCA AGTAGCCAAG TTGGCAGGCA AGTACACAAC AGAAGACGGT TATATCTTTG	1680
	ATCCTCGTGA TATAACCAGT GATGAGGGGG ATGCCTATGT AACTCCACAT ATGACCCATA	1740
	GCCACTGGAT TAAAAAAGAT AGTTTGTCTG AAGCTGAGAG AGCGGCAGCC CAGGCTTATG	1800
45	CTAAAGAGAA AGGTTTGACC CCTCCTTCGA CAGACCACCA GGATTCAGGA AATACTGAGG	1860
	COLOCATION CAGACCACCA GGATTCAGGA AATACTGAGG	1920

17)	·	
	CAAAAGGAGC AGAAGCTATC TACAACCGCG TGAAAGCAGC TAAGAAGGTG	0010000
	STATEGOLIA CAATCITCAA TATACIGTAG AAGTCAAAAA CGGTAGTTTA	3003030
	ATTAIGACCA TTACCATAAC ATCAAATTTG AGTGGTTTGA CGAAGGCCTT	M1001000
	CIAAGGGTA TAGTCTTGAG GATCTTTTGG CGACTGTCAA GTACTATCTC	TATGAGGCAC
	5 ACGAACGTCC GCATTCAGAT AATGGTTTTG GTAACGCTAG TGACCATGTT	GAACATCCAA
	AGGCAGACCA AGATAGTANA CCTGATGAAG ATAAGGAACA TGATGAAGTA	CGTAAAAATA
	CTCACCCTGA ATCTGATGAA AAAGAGAATC ACGCTGGTTT AAATCCTTCA	AGTGAGCCAA
	TTTATAAACC AAGCACTGAT ACGGAAGAGA CAGAGGAAGA AGCTGAAGAT	GCAGATAATC
	AGGCTGAAAT TCCTCAAGTA GAGAATTCTG TTATTAACGC TAAGATAGCA	ACCACAGATG
10	CCTTGCTAGA AAAAGTAACA GATCCTAGTA TTAGACAAAA TGCTATGGAG	GATGCGGAGG
	GTCTAAAAAG TAGTCTTCTT CTCGGAACGA AAGATAATAA CACTATTTCA	ACATTGACTG
	ATAGTCTCTT GGCTTTGTTA AAAGAAAGTC AACCGGCTCC TATACAGTAG	GCAGAAGTAG
	(SEQ ID NO : 5)	TAAAATGAA
	FIGURE 5	
15		
	MKFSKKYIAA GSAVIVSLSL CAYALNQHRS QENKDNNRVS YVDGSQSSQK	
	SENLTPDQVS QKEGIQAEQI VIKITDQGYV TSHGDHYHYY NGKVPYDALF	50
	SEELLMKDPN YOLKDADIVN EVKGGYIIKV DGKYYVYLKD AAHADNVRTK	100
20	DEINRQKQEH VKDNEKVNSN VAVARSQGRY TINDGYVFNP ADIIEDTGNA	150
	YIVPHGGHYH YIPKSDLSAS ELAAAKAHLA GKNMQPSQLS YSSTASDNNT	200
	QSVAKGSTSK PANKSENLQS LLKELYDSPS AQRYSESDGL VFDPAKIISR	250
	TPNGVAIPHG DHYHFIPYSK LSALEEKIAR MVPISGTGST VSTNAKPNEV	300
	VSSLGSLSSN PSSLTTSKEL SSASDGYIFN PKDIVEETAT AYIVRHGDHF	350
25	HYIPKSNQIG QPTLPNNSLA TPSPSLPINP GTSHEKHEED GYGFDANRII	400
	AEDESGFVMS HGDHNHYFFK KDLTEEQIKA AQKHLEEVKT SHNGLDSLSS	450
	HEQDYPGNAK EMKDLDKKIE EKIAGIMKQY GVKRESIVVN KEKNAIIYPH	500
	GDHHHADPID EHKPVGIGHS HSNYELFKPE EGVAKKEGNK VYTGEELTNV	550
	VNLLKNSTFN NQNFTLANGQ KRVSFSFPPE LEKKLGINML VKLITPDGKV	600
30	LEKVSGKVFC EGVGNIANFE LDQPYLPGQT FKYTIASKDY PEVSYDGTFT	650
	VPTSLAYKMA SQTIFYPFHA GDTYLRVNPQ FAVPKGTDAL VRVFDEFHGN	700
	AYLENNYKVG EIKLPIPKLN QGTTRTAGNK IPVTFMANAY LDNQSTYIVE	750
	VPILEKENQT DKPSILPQFK RNKAQENSKL DEKVEEPKTS EKVEKEKLSE	800
	TGNSTSNSTL EEVPTVDPVQ EKVAKFAESY GMKLENVLFN MDGTIELYLP	850
35	SGEVIKKNMA DFTGEAPQGN GENKPSENCK VSTGTVENQP TENKPADSLP	900
	EAPNEKPVKP ENSTDNGMLN PEGNVGSDPM LDPALEEAPA VDPVQEKLEK	950
	FTASYGLGLD SVIFNMDGTI ELRLPSGEVI KKNLSDFIA (SEQ ID NO: 6)	1000
	FIGURE 6	1039
40		
	MKINKKYLAG SVATLVLSVC AYELGLHQAQ TVKENNRVSY IDGKQATQKT	5 0
	ENLTPDEVSK REGINAEQIV IKITDQGYVT SHGDHYHYYN GKVPYDAIIS	50
	EELLMKDPNY QLKDSDIVNE IKGGYVIKVN GKYYVYLKDA AHADNVRTKE	100
	EINRQKQEHS QHREGGTSAN DGAVAFARSQ GRYTTDDGYI FNASDIIEDT	150
45	GDAYIVPHGD HYHYIPKNEL SASELAAAEA FLSGRENLSN LRTYRRQNSD	200
	, TOOLUME DELIKEDED	250

	NTPRINWVPS VSNPGTININ TSNBSNITNS 300	
	NTPRTNWVPS VSNPGTTNTN TSNNSNTNSQ ASQSNDIDSL LKQLYKLPLS 300	
	QRHVESDGLI FDPAQITSRT ARGVAVPHGN HYHFIPYEQM SELEKRIARI 350	
	IPLRYRSNHW VPDSRPEEPS PQPTPEPSPS PQPAPNPQPA PSNPIDEKLV 400	
	KEAVRKVGDG YVFEENGVSR YIPAKNLSAE TAAGIDSKLA KQESLSHKLG AKKTDLPSSD REFYNIKAVDI INDIVIDIO 1888	
	REFINATION LARTHODELD NKGROVDFEA LDNILERIKD 500	
	VSSDKVKLVD DILAFLAPIR HPERLGKPNA QITYTDDEIQ VAKLAGKYTT 550	
	EDGYIFDPRD ITSDEGDAYV TPHMTHSHWI KKDSLSEAER AAAQAYAKEK 600	
	GLTPPSTDHQ DSGNTEAKGA EAIYNRVKAA KKVPLDRMPY NLQYTVEVKN 650	
10	GSLIIPHYDH YHNIKPEWFD EGLYEAPKGY TLEDLLATVK YYVEHPNERP 700	
10	HSDNGFGNAS DHVQRNKNGQ ADTNQTEKPS EEKPQTEKPE EETPREEKPQ 750	
	SERPESPRET EEPEEESPEE SEEPQVETEK VEEKLREAED LLGKIQDPII 800	
	KSNAKETLIG LKNNLLFGTQ DNNTIMAEAE KLLALLKESK (SEQ ID NO: 7) 840	
	PIGURE 7	
15	r	
	MKINKKYLAG SVAVLALSVC SYELGRHQAG QVKKESNRVS YIDGDQAGQK 50	
	AENLTPDEVS KREGINAEQI VIKITDQGYV TSHGDHYHYY NGKVPYDAII 100	
	SEELLMKDPN YQLKDSDIVN EIKGGYVIKV DGKYYVYLKD AAHADNIRTK 150	
	EEIKRQKQEH SHNHNSRADN AVAAARAQGR YTTDDGYIFN ASDIIEDTGD 200	
20	AYIVPHGDHY HYIPKNELSA SELAAAEAYW NGKQGSRPSS SSSYNANPVQ 250	
	PRLSENHNLT VTPTYHQNQG ENISSLLREL YAKPLSERHV ESDGLIFDPA 300	
	QITSRTARGV AVPHGNHYHF IPYEQMSELE KRIARIIPLR YRSNHWVPDS 350	
	RPEQPSPQST PEPSPSLQPA PNPQPAPSNP IDEKLVKEAV RKVGDGYVFE 400	
	ENGVSRYIPA KDLSAETAAG IDSKLAKQES LSHKLGAKKT DLPSSDREFY 450	
25	NKAYDLLARI HQDLLDNKGR QVDFEVLDNL LERLKDVSSD KVKLVDDILA 500	
	FLAPIRHPER LGKPNAQITY TDDEIQVAKL AGKYTTEDGY IFDPRDITSD 550	
	EGDAYVTPHM THSHWIKKDS LSEAERAAAQ AYAKEKGLTP PSTDHQDSGN 600	
	TEAKGAEAIY NRVKAAKKVP LDRMPYNLQY TVEVKNGSLI IPHYDHYHNI 650	
	KFEWFDEGLY EAPKGYSLED LLATVKYYVE HPNERPHSDN GFGNASDHVR 700	
30	KNKADQDSKP DEDKEHDEVS EPTHPESDEK ENHAGLNPSA DNLYKPSTDT 750	
	EETEEEAEDT TDEAEIPQVE NSVINAKIAD AEALLEKVTD PSIRQNAMET 800	
	LTGLKSSLLL GTKDNNTISA EVDSLLALLK ESQPAPIQ 838	
	(SEQ ID NO : 8)	
26	FIGURE 8	
35		
	TGTGCCTATG CACTAAACCA GCATCGTTCG CAGGAAAATA AGGACAATAA TCGTGTCTCT	60
	TATGTGGATG GCAGCCAGTC AAGTCAGAAA AGTGAAAACT TGACACCAGA CCAGGTTAGC	120
40	CAGAAAGAAG GAATTCAGGC TGAGCAAATT GTAATCAAAA TTACAGATCA GGGCTATGTA	180
40	ACGTCACACG GTGATCACTA TCATTACTAT AATGGGAAAG TTCCTTATGA TGCCCTCTTT	240
	AGTGAAGAAC TCTTGATGAA GGATCCAAAC TATCAACTTA AAGACGCTGA TATTGTCAAT	300
	GAAGTCAAGG GTGGTTATAT CATCAAGGTC GATGGAAAAT ATTATGTCTA CCTGAAAGAT	360
	GCAGCTCATG CTGATAATGT TCGAACTAAA GATGAAATCA ATCGTCAAAA ACAAGAACAT	420
15	GICAAAGATA ATGAGAAGGT TAACTCTAAT GTTGCTGTAG CAAGGTCTCA GGGACGATAT	480
45	ACGACAAATG ATGGTTATGT CTTTAATCCA GCTGATATTA TCGAAGATAC GGGTAATGCT	540

•		
	TATATCGTTC CTCATGGAGG TCACTATCAC TACATTCCCA AAAGCGATTT ATCTGCTAGT	
	GAATTAGCAG CAGCTAAAGC ACATCTGGCT GGAAAAAATA TGCAACCGAC TGACTTA	
	INTICTICAL CACCITCICC ATCITCICA ATCAATCCAG GAACTICACA TCACAAACAD	660
	GAAGAAGATG GATACGGATT TGATGCTAAT CGTATTATCG CTGAAGATGA ATCACCTTATT	720
	GICATGAGTC ACGGAGACCA CAATCATTAT TTCTTCAAGA AGGACTTGAC AGAACACCA	780
	ATTAGGCTG CGCAAAAACA TTTAGAGGAA GTTAAAACTA GTCATAATGG ATTAGATTCT	840
	TTGTCATCTC ATGAACAGGA TTATCCAAGT AATGCCAAAG AAATGAAAGA TTTAGATAAA	900
	AAAATCGAAG AAAAAATTGC TGGCATTATG AAACAATATG GTGTCAAACG TGAAAGTATT	960
	GTUGTGAATA AAGAAAAAAA TGCGATTATT TATCCGCATG GAGATCACCA TCATCCACAT	1020
10	CCGATTGATG AACATAAACC GGTTGGAATT GGTCATTCTC ACAGTAACTA TGAACTGTTT	1080
	AAACCCGAAG AAGGAGTTGC TAAAAAAGAA GGGAATAAAG TTTATACTGG AGAAGAATTA	1140
	ACGAATGTTG TTAATTTGTT AAAAAATAGT ACGTTTAATA ATCAAAACTT TACTCTAGCC	1200
	AATGGTCAAA AACGCGTTTC TTTTAGTTTT CCGCCTGAAT TGGAGAAAA ATTAGGTATC	1260
	AATATGCTAG TAAAATTAAT AACACCAGAT GGAAAAGTAT TGGAGAAAGT ATCTGGTAAA	1320
15	GTATTTGGAG AACGAGTAGG GAATATTGCA AACTTTGAAT TAGATCAACC TTATTTACCA	1380
	GGACAACAT TTAAGTATAC TATCGCTTCA AAAGATTATC CAGAAGTAAG TTATGATGGT	1440
	ACATTACAG TTCCAACCTC TTTAGCTTAC AAAATGGCCA GTCAAACGAT TTTCTATCCT	1500
	TTCCATGCAG GGGATACTTA TTTAAGAGTG AACCCTCAAT TTGCAGTGCC TAAAGGAACT	1560
	GATGCTTTAG TCAGAGTGTT TGATGAATTT CATGGAAATG CTTATTTAGA AAATAACTAT	1620
20	AAAGTTGGTG AAATCAAATT ACCGATTCCG AAATTAAACC AAGGAACAAC CAGAACGGCC	1680
	GGAAATAAAA TTCCTGTAAC CTTCATGGCA AATGCTTATT TGGACAATCA ATCGACTTAT	1740
	ATTGTGGAAG TACCTATCTT GGAAAAAGAA AATCAAACTG ATAAACCAAG TATTCTACCA	1800
	CAATTTAAAA GGAATAAAGC ACAAGAAAAC TCAAAACTTG ATGAAAAGGT AGAAGAACCA	1860
	AAGACTAGTG AGAAGGTAGA AAAAGAAAAA CTTTCTGAAA CTGGGAATAG TACTAGTAAT	1920
25	TCAACGTTAG AAGAAGTTCC TACAGTGGAT CCTGTACAAG AAAAAGTAGC AAAATTTGCT	1980
	GAAAGTTATG GGATGAAGCT AGAAAATGTC TTGTTTAATA TGGACGGAAC AATTGAATTA	2040
	TATTTACCAT CGGGAGAAGT CATTAAAAAG AATATGGCAG ATTTTACAGG AGAAGCACCT	2100
	CAAGGAAATG GTGAAAATAA ACCATCTGAA AATGGAAAAG TATCTACTGG AACAGTTGAG	2160
	AACCAACCAA CAGAAAATAA ACCAGCAGAT TCTTTACCAG AGGCACCAAA CGAAAAAACCT	2220
30	GTAAAACCAG AAAACTCAAC GGATAATGGA ATGTTGAATC CAGAAGGGAA TGTGGGGAGT	2280
	GACCCTATGT TAGATTCAGC ATTAGAGGAA GCTCCAGCAG TAGATCCTGT ACAAGAAAAA	2340
	TTAGAAAAAT TTACAGCTAG TTACGGATTA GGCTTAGATA GTGTTATATT CAATATGGAT	2400
	GGAACGATTG AATTAAGATT GCCAAGTGGA GAAGTGATAA AAAAGAATTT ATTGATCTCA	2460
	TAGCGTAA (SEQ ID NO : 9)	2520
35	FIGURE 9	2528
	CAYALNOHRS QENKONNRVS YVDGSQSSQK SENLTPDQVS QKEGIQAEQI 50	
	VIKITDOGYV TSHGDHYUVV NGVUDVDAY B GDDLYAGDAY	
40	EVKGGYIIKV DGKVVVVIKD AAHADAHDDE DEEDONG DEEDONG	
	VAVARSOGRY TTNDCVIEND ADTIBOTON WITHDOWN	
	200 200 200 200 200 200 200 200 200 200	

•	<u> </u>
	TNVVNLLKNS TFNNQNFTLA NGQKRVSFSF PPELEKKLGI NMLVKLITPD 450
	GKVLEKVSGK VFGEGVGNIA NFELDQPYLP GQTFKYTIAS KDYPEVSYDG 500
	TFTVPTSLAY KMASQTIFYP FHAGDTYLRV NPQFAVPKGT DALVRVFDEF 550
_	HGNAYLENNY KVGEIKLPIP KLNQGTTRTA GNKIPVTFMA NAYLDNQSTY 600
5	IVEVPILEKE NOTDKPSILP QFKRNKAQEN SKLDEKVEEP KTSEKVEKEK 650
	LSETGNSTSN STLEEVPTVD PVQEKVAKFA ESYGMKLENV LFNMDGTIEL 700
	YLPSGEVIKK NMADFTGEAP QGNGENKPSE NGKVSTGTVE NQPTENKPAD 750
	SLPEAPNEKP VKPENSTDNG MLNPEGNVGS DPMLDSALEE APAVDPVQEK 800
	LEKFTASYGL GLDSVIFNMD GTIELRLPSG EVIKKNLLIS 840
10	(SEQ ID NO : 10)
	FIGURE 10
. 15	DQGYVTSHGD HYHYYNGKVP YDALFSEELL MKDPNYQLKD ADIVNEVKGG YIIKVDGKYY VYLKDAAHAD NVRTKDEINR QKQEHVKDNE KVNS
	(SEQ ID NO: 11) FIGURE 11
20	GIQAEQIVIK ITDQGYVTSH GDHYHYYNGK VPYDALFSEE LL (SEQ ID NO: 12) FIGURE 12
25	TAYIVRHGDH FHYIPKSNQI GQPTLPNNSL ATPSPSLPI (SEQ ID NO: 13) FIGURE 13
30	TSNSTLEEVP TVDPVQEKVA KFAESYGMKL ENVLFN (SEQ ID NO: 14) FIGURE 14
35	MDGTIELRLP SGEVIKKNLS DFIA (SEQ ID NO: 15) FIGURE 15
40	YGLGLDSVIF NMDGTIELRL PSGEVIKKNL SDFIA (SEQ ID NO: 16) FIGURE 16
	PALEEAPAVD PVQEKLEKFT ASYGLGLDSV IFNMDGTIEL RLPSGEVIKK NLSDFIA (SEQ ID NO: 17)

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FIGURE 17

45

KVEEPKTSEK VEKEKLSETG NSTSNSTLEE VPTVDPVQEK (SEQ ID NO: 18)
FIGURE 18

MKDLDKKIEE KIAGIMKQYG VKRESIVVNK EKNAIIYPHG DHHHADPIDE HKPVGIGHSH
SNYELFKPEE GVAKKEGN
(SEQ ID NO: 19)
FIGURE19

10 AIIYPHGDHH HADPIDEHKP VGICHSHSNY ELFKPEEGVA KKEGNKVYTG E (SEQ ID NO: 20)
FIGURE 20

IQVAKLAGKY TTEDGYIFDP RDITSDEGD (SEQ ID NO: 21)

FIGURE 21

DHQDSGNTEA KGAEAIYNRV KAAKKVPLDR MPYNLQYTVE VKNGSLIIPH YDHYHNIKFE WFDEGLYEAP KGYSLEDLLA TVKYYV

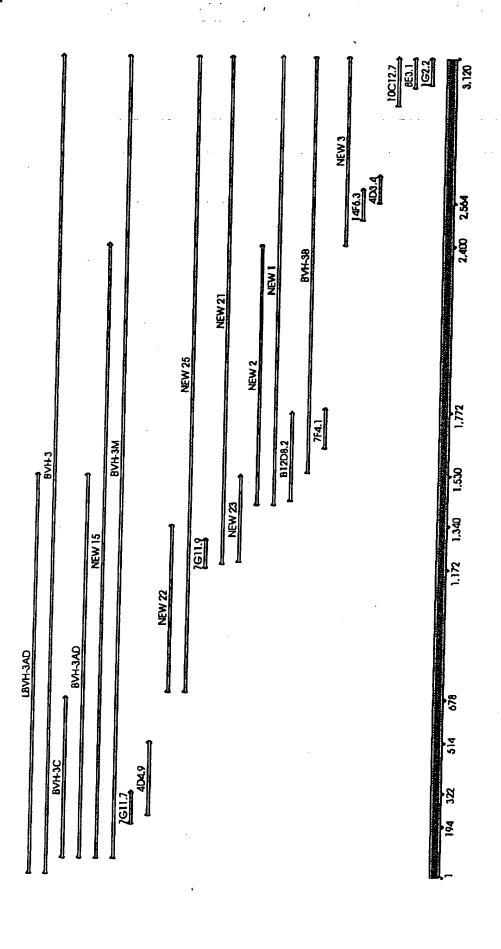
20 (SEQ ID NO: 22) FIGURE 22

GLYEAPKGYS LEDLLATVKY YVEHPNERPH SDNGFGNASD H (SEQ ID NO: 23)

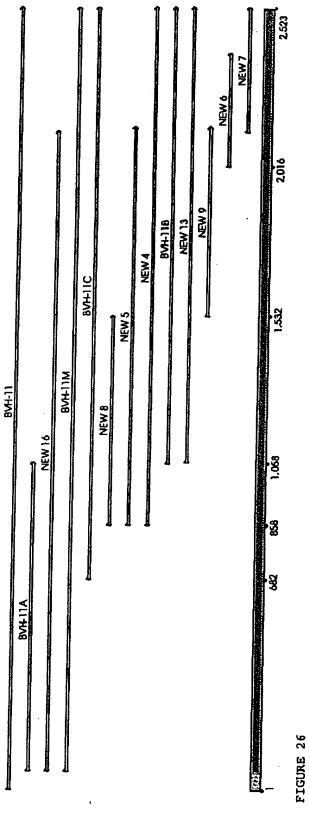
25 FIGURE 23

15

GLYEAPKGYSLEDLLATVKYYV (SEQ ID NO: 163) Figure 24



5 PIGURE 25





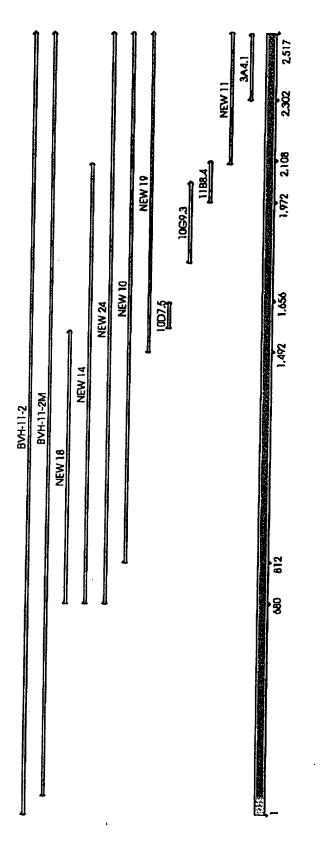
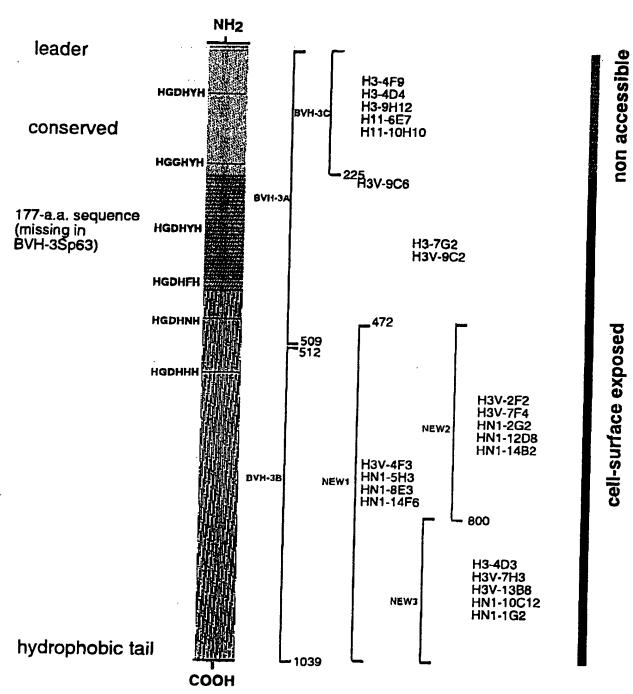
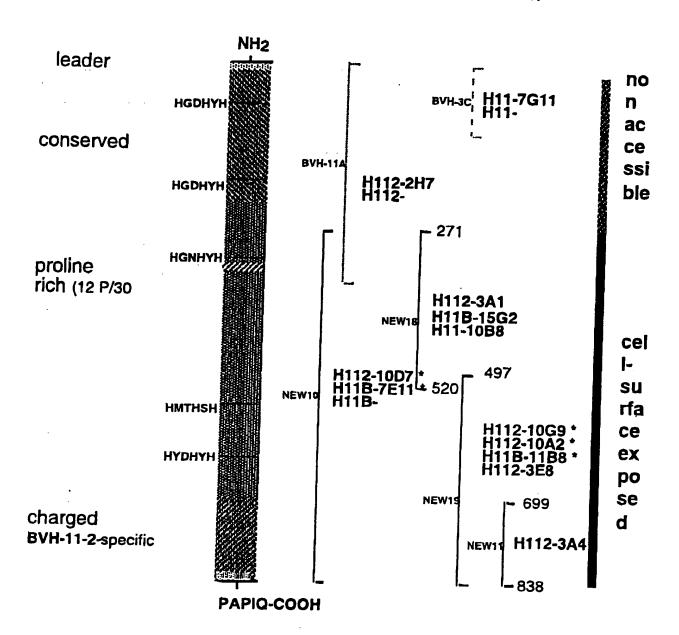


FIGURE 27

Epitope Localization on BVH-3 Protein



Epitope Localization on BVH-11-2 Prot in



^{*} Surface-exposed and protection-conferring Mabs

pury22.His
4925pb

His-tag
Ecorv
Mbal
BamHI
terminator

	BVH-3M	Í	1 CAYALNQHRSQENKDNNRVSYVDGSQSSQKSENLTPDQVSQKEGIQAEQIVIKITDQGYV	
5	5 BVH3-6	3	1 CAYALNQHRSQENKDNNRVSYVDGSQSSQKSENLTPDQVSQKEGIQAEQIVIKITDQGYV	60
				60
	BVH-3M		61 TSHGDHYHYYNGKVPYDALFSEELLMKDPNYQLKDADIVNEVKGGYIIKVDGKYYVYLKD	
	BVH3-6	3	51 TSHGDHYHYYNGKVPYDALFSEELLMKDPNYQLKDADIVNEVKGGYIIKVDGKYYVYLKD	120
10	}		******************************	120
	BVH-3M	1:	1 AAHADNVRTKDEINRQKQEHVKDNEKVNSNVAVARSQGRYTTNDGYVPNPADIIEDTGNA	100
	BVH3-63	3 12	1 AAHADNVRTKDEINROKQEHVKDNEKVNSNVAVARSQGRYTTNDGYVFNPADIIEDTGNA	180
			**************************************	180
15			~~~~~ ~	
	BVH-3M	18	1 YIVPHGGHYHYIPKSDLSASELAAAKAHLAGKNMQPSQLSYSSTASDNNTQSVAKGSTSK	244
	BVH3-63	18	1 YIVPHGGHYHYIPKSDLSASELAAAKAHLAGKOMQPSQLSYSS	240
			******	223
20	BVH-3M	24	1 PANKSENLQSLLKELYDSPSAQRYSESDGLVFDPAKIISRTPNGVAIPHGDHYHFIPYSK	300
	BVH3-63	22	4	300
				223
	BVH-3M	30	LSALEEKIARMVPISGTGSTVSTNAKPNEVVSSLGSLSSNPSSLTTSKELSSASDGYIFN	360
25	BVH3-63	224		223
				223
	,			
	BVH-3M	361	PKDIVEETATAYIVRHGDHFHYIPKSNQIGQPTLPNNSLATPSPSLPINPGTSHEKHEED	420
	BVH3-63	224	TPSPSLPINPGTSHEKHEED	243
30			****	243
	BVH-3M	421	GYGFDANRIIAEDESGFVMSHGDHNHYFFKKDLTEEQIKAAQKHLEEVKTSHNGLDSLSS	480
	BVH3-63	244	GYGFDANRI I A FDESCRIMSHICHLINGUSERVEDI TERCITARIA COMP	303
~-			*********************************	
35				
	BVH-3M	481	HEQDYPGNAKEMKDLDKKIEEKIAGIMKQYGVKRESIVVNKEKNAIIYPHGDHHHADPID	540
	BVH3-63	304	HEQDYPSNAKEMKDLDKKIEEKIAGIMKQYGVKRESIVVNKEKNAIIYPHGDHHHADPID	363

40				
40	BVH-3M	541		500
	BVH3-63	364	EHKPVGIGHSHSNYELFKPEEGVAKKEGNKVYTGEELTNVVNLLKNSTFNNONFTLANGO	123

45	BVH-3M	601	${ t KRVSFSFPPELEKKLGINMLVKLITPDGKVLEKVSGKVFGEGVGNIANFELDQPYLPGQT} = 6$	60
45	BVH3-63	424	KRVSESEPPELEKKI.GINMI VKI IMBDOVAH BIRIGOJA TOTOLOGIA	R3

(4)

	BVH-3M BVH3-63	661 FKYTIASKDYPEVSYDGTFTVPTSLAYKMASQTIFYPFHAGDTYLRVNPQFAVPKGTDAL 720 484 FKYTIASKDYPEVSYDGTFTVPTSLAYKMASQTIFYPFHAGDTYLRVNPQFAVPKGTDAL 543
1(ME-HVE E3-EHVE	721 VRVFDEFHGNAYLENNYKVGEIKLPIPKLNQGTTRTAGNKIPVTFMANAYLDNQSTYIVE 780 544 VRVFDEFHGNAYLENNYKVGEIKLPIPKLNQGTTRTAGNKIPVTFMANAYLDNQSTYIVE 603
	BVH-3M BVH3-63	781 VPILEKENOTOKPSILPOPKRNKAQENSKLDEKVEEPKTSEKVEKEKLSETGNSTSNSTL 840 604 VPILEKENOTOKPSILPOFKRNKAQENSKLDEKVEEPKTSEKVEKEKLSETGNSTSNSTL 663
15	ВVН-3М ВVН3-63	841 EEVPTVDPVQEKVAKFAESYCMKLENVLFNMDGTIELYLPSGEVIKKNMADFTGEAPQGN 900 664 EEVPTVDPVQEKVAKFAESYGMKLENVLFNMDGTIELYLPSGEVIKKNMADFTGEAPQGN 723
20	BVH-3M BVH3-63	901 GENKPSENGKVSTGTVENQPTENKPADSLPEAPNEKPVKPENSTDNGMLNPEGNVGSDPM 960 724 GENKPSENGKVSTGTVENQPTENKPADSLPEAPNEKPVKPENSTDNGMLNPEGNVGSDPM 783
25	BVH-3M BVH3-63 FIGURE 31	961 LDPALEEAPAVDPVQEKLEKFTASYGLGLDSVIFNMDGTIELRLPSGEVIKKNLSDFIA 1019 784 LDSALEEAPAVDPVQEKLEKFTASYGLGLDSVIFNMDGTIELRLPSGEVIKKNLLIS 840
30		
35	BVH-3 BVH-11 BVH-11-2	1 MKFSKKYIAAGSAVIVSLSLCAYALNQHRSQENK-DNNRVSYVDGSQSSQKSENLTPDQV 59 1 MKINKKYLAG-SVATLVLSVCAYELGLHQAQTVK-ENNRVSYIDGKQATQKTENLTPDEV 58 1 MKINKKYLAG-SVAVLALSVCSYELGRHQAGQVKKESNRVSYIDGDQAGQKAENLTPDEV 59 ** *** * * * * * * * * * * * * * * * *
40	BVH-3 BVH-11 BVH-11-2	60 SQKEGIQAEQIVIKITDQGYVTSHGDHYHYYNGKVPYDALFSEELLMKDPNYQLKDADIV 119 59 SKREGINAEQIVIKITDQGYVTSHGDHYHYYNGKVPYDAIISEELLMKDPNYQLKDSDIV 118 60 SKREGINAEQIVIKITDQGYVTSHGDHYHYYNGKVPYDAIISEELLMKDPNYQLKDSDIV 119 **.***
45	BVH-3 BVH-11 BVH-11-2	120 NEVKGGYIIKVDGKYYVYLKDAAHADNVRTKDEINRQKQEHVKDNEKVNSNVAVAR 175 119 NEIKGGYVIKVNGKYYVYLKDAAHADNVRTKEEINRQKQEHSQHREGGTSANDGAVAFAR 178 120 NEIKGGYVIKVDGKYYVYLKDAAHADNIRTKEEIKRQKQEHSHNHNSRADNAVAAAR 176

	BVH-3	176 SOGRYTTNDGVIENDADITETTOTAL
	BVH-11	176 SQGRYTTNDGYVFNPADIIEDTGNAYIVPHGGHYHYIPKSDLSASELAAAKAHLAGKNMQ 235
	BVH-11-2	179 SQGRYTTDDGYIFNASDIIEDTGDAYIVPHGDHYHYIPKNELSASELAAAEAFLSGRENL 238
		2 177 AQGRYTTDDGYIFNASDIIEDTGDAYIVPHGDHYHYIPKNELSASELAAAEAYWNGKQ 234
		* **************
	BVH-3	236 PSQLSYSSTASDNNTQSVAKGSTSKPA-NKSENLQSLLKELYDSP 279
	BVH-11	239 SNLRTYRRONSDNTPRTNWVPSVSNPCTTNTNTSDNICNTNICO COGNATION
	BVH-11-2	235 -GSRPSSSSYNANPVOPRI.SENIMI.TETEPTENIA
10	•	*
	BVH-3	280 SAQRYSESDGLVFDPAKIISRTPNGVAIPHGDHYHFIPYSKLSALEEKIARMVPISGTGS 339
	BVH-11	299 LSQRHVESDGLIFDPAQITSRTARGVAVPHGNHYHFIPYEQMSELEKRIARIIPLRYRSN 358
	BVH-11-2	285 LSERHVESDGLIFDPAQITSRTARGVAVPHGNHYHFIPYEQMSELEKRIARIIPLRYRSN 344
15		* * **** *** * *** *** *** *** ***
	BVH-3	340 TVSTNAKPNEVVSSLGSLSSNPSSLTTSKELSSASDGYIFNPKDIVEETATAYIVRHGDH 399
	BVH-11	359 HWVPDSRP-EEPSPQPTPEPSPS-PQPAPNPQPAPSNPIDEKLVKEAVRKVGDG 410
20	BVH-11-2	345 HWVPDSRP-EQPSPQSTPEPSPS-LQPAPNPQPAPSNPIDEKLVKEAVRKVGDG 396
20		
	BVH-3	400 FHYTPKSNOTGOPTI DADIEL AMDEREL DENGE CHIEF
	BVH-11	400 FHYIPKSNQIGQPTLPNNSLATPSPSLPINPGTSHEKHEEDGYGFDANRIIAEDESGFVM 459 411 YVFEENGVSRYIPAKNLSAETAAGIDSKLAKQESLS 446
	BVH-11-2	397 YVFEENCVSPVID AVDY CARDAL CONTRACTOR
25		* . * * * * * * * * * * * * * * * * * *
		•••
	BVH-3	460 SHGDHNHYFFKKDLTEEQIKAAQKHLEEVKTSHNGLDSLSSHEQDYPGNAKEMKDLDKKI 519
	BVH-11	447HKLGAKKTDLPSSDREFYNKAYDLLARIHQDLLDNKGRQVDFEALDNLLERLKDVS 502
	BVH-11-2	433HKLGAKKTDLPSSDREFYNKAYDLLARIHQDLLDNKGRQVDFEVLDNLLERLKDVS 488
30		* * * * * * * * * * * * * * * * * * * *
	BVH-3	520 EEKIAGIMKQYGVKRESIVVNKEKNAIIYPHGDHHHADPIDEHKPVGIGHSHSNYELFKP 579
	BVH-11 BVH-11-2	503 SDKVKLVDDILAFLAPIRHPERLGKPNAQITYTDDEIQVAKLAGKYTTEDGYIFDP 558
35	BVN-11-2	489 SDKVKLVDDILAFLAPIRHPERLGKPNAQITYTDDEIQVAKLAGKYTTEDGYIFDP 544
00		
	BVH-3	580 EEGVAKKEGNIGIVYGEET THRADII I VALGODADAN AND AND AND AND AND AND AND AND AN
	BVH-11	580 EEGVAKKEGNKVYTGEELTNVVNLLKNSTFNNQNFTLANGQKRVSFSPPPELEKKLGINM 639 559 RD-ITSDEGD-AYVTPHMTHSHWIKKDS-LSEAERAAAQAYAKEKGLTPPSTDHQD 611
	BVH-11-2	545 RD-ITSDEGD-AYVTPHMTHSHWIKKDS-LSEAERAAAQAYAKEKGLTPPSTDHQD 597
40		•• • • • • • • • • • • • • • • • • • • •
•		
	BVH-3	640 LVKLITPDGKVLEKVSGKVFGEGVGNIANFELDQPYLPGQTFKYTIASKDYPEVSYDGTF 699
	BVH-11	612SGNTEAKGAEAIYNRVKAAKKVPLDRMPYNLQYTVEVKNGSL 653
	BVH-11-2	598SGNTEAKGAEAIYNRVKAAKKVPLDRMPYNLQYTVEVKNGSL 639
45		** * * * * * * * * * * * * * * * * * * *

	BVH-3	700 TVPTSLAYKMASOTTEVPEURODERS TIPE	
	BVH-11	700 TVPTSLAYKMASQTIFYPFHAGDTYLRVNPQPAVPKGTDALVRVFDEFHGNAYLENNYKV 75	9
	BVH-11-2	654 IIPHYDHYHNIKFEWFDEG	8
5		2 640 IIPHYDHYHNIKFEWFDEGLYEAPKGYSLEDLLAT 67	4
_		**	-
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		· · · · · · · · · · · · · · · · · · ·	
1.	BVH-3	760 GEIKLPIPKLNQGTTRTAGNKIPVTFMANAYLDNQSTYIVEVPILEKENQTDKPSILPQP 819	
10		TO STATE OF THE PROPERTY OF TH	
	BVH-11-2	6/5 VKYYVEHPNERPHSDNGFGNASDHIJPW	-
		710)
		• • • •	
	BVH-3	820 KRNKAQENSKLDEKVEEPKTSEKVEKEKLSETGNSTSNSTLEEVPTVDPVQEKVAKFAES 879	
15	BVH-11	725QTEKPSEEKPQTEKPEEE	
	BVH-11-2	711DEDKEHDEVSEPTHPESDEKE	
		* * * * 731	
		* * *	
	BVH-3	880 VOWE ENDINGERS IN THE STATE OF THE STATE	
20	BVH-11	880 YGMKLENVLFNMDGTIELYLPSGEVIKKNMADFTGEAPQGNGENKPSENGKVSTGTVENQ 939	
	BVH-11-2	743	
	BVN-11-2	732TE 751	
		* **	
25			
23	BVH-3	940 PTENKPADSLPEAPNEKPVKPENSTDNGMLNPEGNVGSDPMLDPALEEAPAVDPVQEKLE 999	
	BVH-11	759 PTEEPEESPEESEEPQVETEKVEEKLREAEDLLGKIODPIIKSNAKETLT ROD	
	BVH-11-2	752 ETEEEAEDTTDEAEIPQVENSVINAKIADAEALLEKVTDPSIRQNAMETLT 802	
		** * * * * * * * * * * * * * * * * * * *	
30	BVH-3	1000 KFTASYGLGLDSVIFNMDGTIELRLPSGEVIKKNLSDFIA 1039	
	BVH-11	810 GLKNNT.PGTO- DARMING BARRES	
	BVH-11-2	803 GLKSSLLLCTV	
		* **	
		• • • • • • • • • • • • • • • • • • •	
35	FIGURE 32		

5 10 15	61 ACAGAAGACG CTATATCTT TGATACTAGT TGGATTAAAA AAGATAGTTT GTCTGAAGCT 121 GAGAGAGCG CTATATCTT TGATACTAGT 181 CACCAGGATT CAGGAAATAC TGAGGCAAAA GAGACAGAGGTT CAGCAGGATT CAGGAAATAC TGAGGCAAAA GGAGCAGAAG CTATCTACAA CCGCGTGAAA 301 AAAAACGGTA GTTTAATCAT ACCTCATTAT GACCATTACC ATAACATCAA ATTTGAGTAG 361 TTTGACGAAG GCCTTTATGA GGCACCTAAG GGGTATAGTC TTGAGGATCT TTTGAGGAGTC 421 GTCAAGTACT ATGTCGAACC GCGGAACGCT AGTGACCATG TTCGTAAAAA ATTTGAGTAG 421 GAATCTGATGA AACCTGATGA AGATAAGGAA CATGATGAAG TAAGTGAGCC AACTCACCCT 481 CAAGATAGTA AACCTGATGA AGATAAGGAA CATGATGAAG TAAGTGAGCC AACTCACCCT 601 CCAAGCACTG ATACGGAAGA GAAGAGGAA GAAGCTGAAG TAAGTGAGCC CAGCAGATAA TCTTTATAAA 661 ATTCCTGGTA CCCCTAGTAT TAGACAAAAT GCTATGAGA CATTGACTGG TCTAAAAAGT 721 AGTCTTCTTC TCGGAACGAA ACCGGCTCCT ATACAGTAG (SEQ ID NO: 257)
20	FIGURE 33
25	1 MQITYTDDEI QVAKLAGKYT TEDGYIFDTS WIKKDSLSEA ERAAAQAYAK EKGLTPPSTD 61 HQDSGNTEAK GAEAIYNRVK AAKKVPLDRM PYNLQYTVEV KNGSLIIPHY DHYHNIKFEW 121 FDEGLYEAPK GYSLEDLLAT VKYYVEPRNA SDHVRKNKAD QDSKPDEDKE HDEVSEPTHP 181 ESDEKENHAG LNPSADNLYK PSTDTEETEE EAEDTTDEAE IPGTPSIRQN AMETLTGLKS 241 SLLLGTKDNN TISAEVDSLL ALLKESQPAP IQ (SEQ ID NO : 258)
.JU	